SEQUENCE LISTING

<110> Mize, Nancy K. Haley-Vicente, Dana A.
<120> Interleukin-1 Hy2 Materials and Methods
<130> 28110/36858A
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<150> US 60/245,346 <151> 2000-11-02
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Cys Ser Leu Pro Met Ala Arg Tyr Tyr Ile Ile Lys Tyr Ala Asp Gln
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Lys Ala Leu Tyr Thr Arg Asp Gly Gln Leu Leu Val Gly Asp Pro Val
20 25 30

HONOTO EN HE INCIDENTAL 15 TO BE

gca	gac	aac	tgc	tgt	gca	gag	aag	atc	tgc	aca	ctt	cct	aac	aga	ggc	200
Ala	Asp	Asn	Cys	Cys	Ala	Glu	Lys	Ile	Cys	Thr	Leu	Pro	Asn	Arg	Gly	
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Leu	Asp	Arg	Thr	Lys	Val	Pro	Ile	Phe	Leu	Gly	Ile	Gln	Gly	Gly	Ser	
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Arg	Cys	Leu	Ala	Cys	Val	Glu	Thr	Glu	Glu	Gly	Pro	Ser	Leu	Gln	Leu	
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Glu	Asp	Val	Asn	Ile	Glu	Glu	Leu	Tyr	Lys	Gly	Gly	Glu	Glu	Ala	Thr	
			85					90					95			
999	++0	200	++0	++-	<i>a</i> 2 <i>a</i>	200	200	tas	~~~	+	~~~	++-	200	a++	~~~	392
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Arg	Pile		Pne	Pne	GIII	261		ser	GIÀ	ser	Ald		Arg	ьeu	GIU	
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	115					120					125					
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Gln	Pro	Val	Gln	Leu	Thr	Lys	Glu	Ser	Glu	Pro	Ser	Ala	Arg	Thr	Lys	
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Phe	Tyr	Phe	Glu	Gln	Ser	Trp										

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40

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Gly Leu Asp Arg Thr Lys Val Pro Ile Phe Leu Gly Ile Gln Gly Gly
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Ser Arg Cys Leu Ala Cys Val Glu Thr Glu Glu Gly Pro Ser Leu Gln 65 70 75 80

Leu Glu Asp Val Asn Ile Glu Glu Leu Tyr Lys Gly Glu Glu Ala 85 90 95

Thr Arg Phe Thr Phe Phe Gln Ser Ser Ser Gly Ser Ala Phe Arg Leu
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Glu Ala Ala Trp Pro Gly Trp Phe Leu Cys Gly Pro Ala Glu Pro 115 120 125

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35 40 45

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50 55 60

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Leu	Gln	Leu	Glu	Asp	Val	Asn	Ile	Glu	Glu	Leu	Tyr	Lys	Gly	Gly	Glu	
				100					105					110		
gag	acc	aca	cac	++ <i>c</i>	acc	ttc	ttc	cad	add	add	tca	aac	taa	acc	ttc	383
Glu			_													-
-	-		115					120				1	125			
agg	ctt	gag	gct	gct	gcc	tgg	cct	ggc	tgg	ttc	ctg	tgt	ggc	ccg	gca	431
Arg	Leu	Glu	Ala	Ala	Ala	Trp	Pro	Gly	Trp	Phe	Leu	Cys	Gly	Pro	Ala	
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Glu		Gln	Gln	Pro	Val		Leu	Thr	Lys	Glu		Glu	Pro	Ser	Ala	
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Arg											33	_				
160					165					170						
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20 25 30

Asp Gln Lys Ala Leu Tyr Thr Arg Asp Gly Gln Leu Leu Val Gly Asp

45

Pro Val Ala Asp Asn Cys Cys Ala Glu Lys Ile Cys Thr Leu Pro Asn 50 55 60

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155

160

Arg Gly Leu Asp Arg Thr Lys Val Pro Ile Phe Leu Gly Ile Gln Gly 65 70 75 80 Gly Ser Arg Cys Leu Ala Cys Val Glu Thr Glu Glu Gly Pro Ser Leu 90 95 85 Gln Leu Glu Asp Val Asn Ile Glu Glu Leu Tyr Lys Gly Gly Glu Glu 100 105 110 Ala Thr Arg Phe Thr Phe Phe Gln Ser Ser Ser Gly Ser Ala Phe Arg 120 125 115 Leu Glu Ala Ala Trp Pro Gly Trp Phe Leu Cys Gly Pro Ala Glu 130 135 140 Pro Gln Gln Pro Val Gln Leu Thr Lys Glu Ser Glu Pro Ser Ala Arg

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20 25 30

Ala Gly Lys Val Ile Lys Gly Glu Glu Ile Ser Val Val Pro Asn Arg

40

45

Trp Leu Asp Ala Ser Leu Ser Pro Val Ile Leu Gly Val Gln Gly Gly
50 55 60

Ser Gln Cys Leu Ser Cys Gly Val Gly Gln Glu Pro Thr Leu Thr Leu 65 70 75 80

Glu Pro Val Asn Ile Met Glu Leu Tyr Leu Gly Ala Lys Glu Ser Lys

85 90 95

Ser Phe Thr Phe Tyr Arg Arg Asp Met Gly Leu Thr Ser Ser Phe Glu
100 105 110

Ser Ala Ala Tyr Pro Gly Trp Phe Leu Cys Thr Val Pro Glu Ala Asp 115 120 125

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Pro Ile Thr Asp Phe Tyr Phe Gln Gln Cys Asp
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<211> 178

<212> PRT

<213> Rattus rattus

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5

10

15

Ile Leu Leu Phe Arg Ser Glu Ser Ala Gly His Pro Ala Gly Lys Arg

20

25

30

Pro Cys Lys Met Gln Ala Phe Arg Ile Trp Asp Thr Asn Gln Lys Thr

35

40

45

Phe Tyr Leu Arg Asn Asn Gln Leu Ile Ala Gly Tyr Leu Gln Gly Pro

50 55 6

Asn Thr Lys Leu Glu Glu Lys Ile Asp Met Val Pro Ile Asp Phe Arg
65 70 75 80

Asn Val Phe Leu Gly Ile His Gly Gly Lys Leu Cys Leu Ser Cys Val

Lys Ser Gly Asp Asp Thr Lys Leu Gln Leu Glu Glu Val Asn Ile Thr

Asp Leu Asn Lys Asn Lys Glu Glu Asp Lys Arg Phe Thr Phe Ile Arg

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Asn Thr Pro Lys Glu Pro Cys Thr Val Thr Lys Phe Tyr Phe Gln Glu

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Asp Gln

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35 40 45

Tyr Leu Arg Asn Asn Gln Leu Val Ala Gly Tyr Leu Gln Gly Pro Asn
50 55 60

Thr Lys Leu Glu Glu Lys Ile Asp Val Val Pro Val Glu Pro His Phe
65 70 75 80

Val Phe Leu Gly Ile His Gly Gly Lys Leu Cys Leu Ser Cys Val Lys

85 90 95

Ser Gly Asp Glu Met Lys Leu Gln Leu Asp Ala Val Asn Ile Thr Asp

100 105 110

Leu Arg Lys Asn Ser Glu Gln Asp Lys Arg Phe Thr Phe Ile Arg Ser

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Tyr Leu Arg Asn Asn Gln Leu Val Ala Gly Tyr Leu Gln Gly Pro Asn
50 55 60

Val Asn Leu Glu Glu Lys Ile Asp Val Val Pro Ile Glu Pro His Ala
65 70 75 80

Leu Phe Leu Gly Ile His Gly Gly Lys Met Cys Leu Ser Cys Val Lys

85 90 95

Ser Gly Asp Glu Thr Arg Leu Gln Leu Glu Ala Val Asn Ile Thr Asp

100 105 110

Leu Ser Glu Asn Arg Lys Gln Asp Lys Arg Phe Ala Phe Ile Arg Ser 115 120 125

Asp Ser Gly Pro Thr Thr Ser Phe Glu Ser Ala Ala Cys Pro Gly Trp

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Glu

<210> 9

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<212> PRT

<213> Homo sapiens

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Pro	Ser	Leu	Leu	Pro	Ile	ser	Glu	Asp	Gln	Thr	Pro	Leu	Ile	Ala	Gly
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Lys Phe Tyr Phe Glu Gln Ser Trp

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<212> DNA

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THE HARLEST THE REPORT OF THE

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<213> Homo sapiens

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<210> 16

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<212> DNA

<213> Mouse

<400> 16

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<210> 17

<211> 459

<212> DNA

<213> Mouse

<400> 17

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<210> 18

<211> 152

<212> PRT

<213> Mouse

<400> 18

Met Cys Ser Leu Pro Met Ala Arg Tyr Tyr Ile Ile Lys Asp Ala His 1 5 10 15

Gln Lys Ala Leu Tyr Thr Arg Asn Gly Gln Leu Leu Gly Asp Pro 20 25 30

Asp Ser Asp Asn Tyr Ser Pro Glu Lys Val Cys Ile Leu Pro Asn Arg

40

45

Gly Leu Asp Arg Ser Lys Val Pro Ile Phe Leu Gly Met Gln Gly Gly
50 55 60

Ser Cys Cys Leu Ala Cys Val Lys Thr Arg Glu Gly Pro Leu Leu Gln
65 Val Asp Val Asn Ile Glu Asp Leu Tyr Lys Gly Gly Glu Gln Thr

Led Gid Asp val Ash lie Gid Asp Led lyl Lys Cly Cly Cld Cld 1911

<220>

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<223> Description of Artificial Sequence: primer

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Cys Thr Ala Met Glu Ala Asp Gln Pro Val Ser Leu Thr Asn Met Pro

Asp Glu Gly Val Met Val Thr Lys Phe Tyr Phe Gln Glu Asp Glu

130 135 140

<210> 22

<211> 146

<212> PRT

<213> Homo sapiens

<400> 22

Ala Arg Tyr Tyr Ile Ile Lys Tyr Ala Asp Gln Lys Ala Leu Tyr Thr

1 5 10 15

Arg Asp Gly Gln Leu Leu Val Gly Asp Pro Val Ala Asp Asn Cys Cys

20 25 30

Ala Glu Lys Ile Cys Ile Leu Pro Asn Arg Gly Leu Asp Arg Thr Lys

35 40 45

Val Pro Ile Phe Leu Gly Ile Gln Gly Gly Ser Arg Cys Leu Ala Cys

50 55 60

Val Glu Thr Glu Glu Gly Pro Ser Leu Gln Leu Glu Asp Val Asn Ile

65 70 75 80

Glu Glu Leu Tyr Lys Gly Glu Glu Glu Ala Thr Arg Phe Thr Phe Phe

90 95

Gln Ser Ser Ser Gly Ser Ala Phe Arg Leu Glu Ala Ala Ala Trp Pro

100 105 110

Gly Trp Phe Leu Cys Gly Pro Ala Glu Pro Gln Gln Pro Val Gln Leu

115 120 125

ייי וואסקות אינו אין אוויי או לייינו אין אוויי אוו

Thr Lys Glu Ser Glu Pro Ser Ala Arg Thr Lys Phe Tyr Phe Glu Gln
130 135 140

Ser Trp

145

<210> 23

<211> 151

<212> PRT

<213> Homo sapiens

<400> 23

Val Arg Ser Leu Asn Cys Thr Leu Arg Asp Ser Gln Gln Lys Ser Leu

1 5 10 15

Val Met Ser Gly Pro Tyr Glu Leu Lys Ala Leu His Leu Gln Gly Gln
20 25 30

Asp Met Glu Gln Gln Val Val Phe Ser Met Ser Phe Val Gln Gly Glu

35 40 45

Glu Ser Asn Asp Lys Ile Pro Val Ala Leu Gly Leu Lys Glu Lys Asn 50 55 60

Leu Tyr Leu Ser Cys Val Leu Lys Asp Asp Lys Pro Thr Leu Gln Leu 65 70 75 80

Glu Ser Val Asp Pro Lys Asn Tyr Pro Lys Lys Met Glu Lys Arg
85 90 95

Phe Val Phe Asn Lys Ile Glu Ile Asn Asn Lys Leu Glu Phe Glu Ser

Ala Gln Phe Pro Asn Trp Tyr Ile Ser Thr Ser Gln Ala Glu Asn Met
115 120 125

Pro Val Phe Leu Gly Gly Thr Lys Gly Gly Gln Asp Ile Thr Asp Phe
130 135 140

Thr Met Gln Phe Val Ser Ser

145 150

<210> 24

<211> 148

<212> PRT

<213> Homo sapiens

<400> 24

Pro Met Ala Arg Tyr Tyr Ile Ile Lys Tyr Ala Asp Gln Lys Ala Leu

1 5 10 15

Tyr Thr Arg Asp Gly Gln Leu Leu Val Gly Asp Pro Val Ala Asp Asn
20 25 30

Cys Cys Ala Glu Lys Ile Cys Ile Leu Pro Asn Arg Gly Leu Asp Arg

40

45

Thr Lys Val Pro Ile Phe Leu Gly Ile Gln Gly Gly Ser Arg Cys Leu 50 55 60

Ala Cys Val Glu Thr Glu Glu Gly Pro Ser Leu Gln Leu Glu Asp Val 80 70 75 65 Asn Ile Glu Glu Leu Tyr Lys Gly Glu Glu Ala Thr Arg Phe Thr 95 85 90 Phe Phe Gln Ser Ser Ser Gly Ser Ala Phe Arg Leu Glu Ala Ala Ala 100 105 110 Trp Pro Gly Trp Phe Leu Cys Gly Pro Ala Glu Pro Gln Gln Pro Val 115 120 125 Gln Leu Thr Lys Glu Ser Glu Pro Ser Ala Arg Thr Lys Phe Tyr Phe 130 135 140 Glu Gln Ser Trp 145 <210> 25 <211> 28 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Primer <400> 25 28 gtcatatgtg ttccctcccc atggcaag

<210> 2

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 26

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37